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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/784,870	02/24/2004	Mikio Takaiwa	249142US0DIV	1309
22850	7590	06/12/2006	EXAMINER	
OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT, P.C. 1940 DUKE STREET ALEXANDRIA, VA 22314			RAO, MANJUNATH N	
			ART UNIT	PAPER NUMBER
			1652	
DATE MAILED: 06/12/2006				

Please find below and/or attached an Office communication concerning this application or proceeding.

Office Action Summary	Application No. 10/784,870	Applicant(s) TAKAIWA ET AL.	
	Examiner Manjunath N. Rao, Ph.D.	Art Unit 1652	

-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --

Period for Reply

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) OR THIRTY (30) DAYS, WHICHEVER IS LONGER, FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

Status

- 1) ☒ Responsive to communication(s) filed on 05 December 2005.
- 2a) ☐ This action is FINAL. 2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

Disposition of Claims

- 4) ☒ Claim(s) 7-34 is/are pending in the application.
- 4a) Of the above claim(s) _____ is/are withdrawn from consideration.
- 5) ☐ Claim(s) _____ is/are allowed.
- 6) ☒ Claim(s) 7-34 is/are rejected.
- 7) ☐ Claim(s) _____ is/are objected to.
- 8) ☐ Claim(s) _____ are subject to restriction and/or election requirement.

Application Papers

- 9) ☒ The specification is objected to by the Examiner.
- 10) ☒ The drawing(s) filed on 24 February 2004 is/are: a) ☒ accepted or b) ☐ objected to by the Examiner.
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

Priority under 35 U.S.C. § 119

- 12) ☒ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☒ All b) ☐ Some * c) ☐ None of:
1. ☐ Certified copies of the priority documents have been received.
 2. ☒ Certified copies of the priority documents have been received in Application No. 09/920,954.
 3. ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).

* See the attached detailed Office action for a list of the certified copies not received.

Attachment(s)

- | | |
|---|---|
| 1) <input checked="" type="checkbox"/> Notice of References Cited (PTO-892) | 4) <input type="checkbox"/> Interview Summary (PTO-413)
Paper No(s)/Mail Date. _____ |
| 2) <input type="checkbox"/> Notice of Draftsperson's Patent Drawing Review (PTO-948) | 5) <input type="checkbox"/> Notice of Informal Patent Application (PTO-152) |
| 3) <input checked="" type="checkbox"/> Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08)
Paper No(s)/Mail Date <u>2/04</u> . | 6) <input checked="" type="checkbox"/> Other: <u>sequence alignments(2 p)</u> . |

DETAILED ACTION

Claims 7-34 are currently pending in this application.

Drawings

Drawings submitted in this application are accepted by the Examiner for examination purposes only.

Specification

Examiner notes that applicants have not updated the relationship of the instant application to its parent application that has matured in to a US patent. Examiner urges applicants to amend said information by providing the US patent number in response to this Office action.

Claim Rejections - 35 USC § 112

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 18 and 32 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention. Claims 18 and 32 are drawn to the same biological deposits. However each claim claims that the polynucleotide in these biological deposit are different. For example the biological deposit of claim 18 indicates it comprises the polynucleotide which encodes the polypeptide with SEQ ID NO:1 which that of claim 32, claims a deposit comprising

Art Unit: 1652

polynucleotide encoding SEQ ID NO:2. It is not clear to the Examiner as to how a single deposit can comprise two different polynucleotides. Examiner requests clarification.

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

Claims 18 and 32 are rejected because the invention appears to employ microorganisms transformed with novel polynucleotides. Since the microorganisms are essential to the claimed invention, they must be obtainable by a repeatable method set forth in the specification or otherwise be readily available to the public. The claimed microorganisms' sequences are not fully disclosed, nor have all the sequences required for their construction been shown to be publicly known and freely available. The specification does not disclose a repeatable process to obtain the microorganisms and it is not apparent if the DNA sequences are readily available to the public. Accordingly, it is deemed that a deposit of these microorganisms should have been made in accordance with 37 CFR 1.801-1.809. In order for the claims to be enabled, applicants must show that either the microorganisms can be made by publicly available materials or that the microorganisms as such has been deposited in such a way that it is freely available to the public. The enablement requirements of 35 U.S.C. § 112 may be satisfied by a deposit of the plasmids and the host cells that are transformed using said plasmids.

It appears that applicants have made a deposit under the terms of the Budapest Treaty. Therefore an affidavit or declaration by applicants, or a statement by an attorney of record over his or her signature and registration number, stating that the specific plasmid/strain has been

Art Unit: 1652

deposited under the Budapest Treaty and that the strain will be irrevocably and without restriction or condition released to the public upon the issuance of the patent, would satisfy the deposit requirement made herein.

Claims 7-34 are rejected under 35 U.S.C. 112, first paragraph, because the specification, while being enabling for a polynucleotide encoding a polypeptide with SEQ ID NO:1 or 2 and having a specific protease activity, vectors and host cells comprising said polynucleotide and a method of making the polypeptide using the host cell comprising said polynucleotide, does not reasonably provide enablement for any such polynucleotide which encodes a polypeptide that is at least 90% identical to SEQ ID NO:1 or 2 and encoding a polypeptide having protease activity, vectors and host cells comprising said polynucleotide and a method of making the polypeptide using the host cell comprising said polynucleotide. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make the invention commensurate in scope with these claims.

Factors to be considered in determining whether undue experimentation is required, are summarized in *In re Wands* (858 F.2d 731, 8 USPQ 2nd 1400 (Fed. Cir. 1988)) as follows: (1) the quantity of experimentation necessary, (2) the amount of direction or guidance presented, (3) the presence or absence of working examples, (4) the nature of the invention, (5) the state of the prior art, (6) the relative skill of those in the art, (7) the predictability or unpredictability of the art, and (8) the breadth of the claim(s).

Claims 7-34 are so broad as to encompass any polynucleotide that encodes a polypeptide that is at least 90% identical in its sequence to SEQ ID NO:1 or 2 vectors and host cells

Art Unit: 1652

comprising said polynucleotide and a method of making the polypeptide using the host cell comprising said polynucleotide. The scope of the claims is not commensurate with the enablement provided by the disclosure with regard to the extremely large number of polynucleotides broadly encompassed by the claims. Since the amino acid sequence of a protein encoded by said polynucleotide determines its structural and functional properties, predictability of which changes can be tolerated in said encoded protein's amino acid sequence and obtain the desired activity requires a knowledge of and guidance with regard to which amino acids in the protein's sequence, if any, are tolerant of modification and which are conserved (i.e. expectedly intolerant to modification), and detailed knowledge of the ways in which the proteins' structure relates to its function. Simply put, applicants have not taught those skilled in the art as to where exactly on the polynucleotide sequence encoding SEQ ID NO:1 or 2, specific nucleotides can be modified (i.e., by insertion, deletion or substitution), and how to select those modified sequences in order to arrive at those that encode the polypeptide having the specific activity of SEQ ID NO:1 or 2. The specification is limited to teaching the use of the polynucleotide with SEQ ID NO:3 or 4 to encode the polypeptide with SEQ ID NO:1 or 2 and use it as a specific protease but provides no guidance with regard to the making of variants and mutants or with regard to the other uses indicated above. In view of the great breadth of the claim, amount of experimentation required to make the claimed polypeptides, the lack of guidance, working examples, and unpredictability of the art in predicting function from a polypeptide primary structure (e.g., see Ngo et al. in *The Protein Folding Problem and Tertiary Structure Prediction*, 1994, Merz et al. (ed.), Birkhauser, Boston, MA, pp. 433 and 492-495, Ref: U, Form-892), the claimed invention

Art Unit: 1652

would require undue experimentation. As such, the specification fails to teach one of ordinary skill how to use the full scope of the polypeptides encompassed by this claim.

While recombinant and mutagenesis techniques are known, it is not routine in the art to screen for multiple substitutions or multiple modifications, as encompassed by the instant claims, and the positions within a polynucleotide sequence leading to variants or mutants through which amino acid modifications can be made with a reasonable expectation of success in obtaining the desired activity/utility are limited in any encoded protein, and the result of such modifications is unpredictable. In addition, one skilled in the art would expect any tolerance to modification for a given protein to diminish with each further and additional modification, e.g. multiple substitutions.

The specification does not support the broad scope of the claims which encompass all modifications and fragments of any polynucleotide encoding a polypeptide with 90% sequence identity to SEQ ID NOS:1 or 2 because the specification does not establish: (A) regions of the polynucleotide structure which may be modified without affecting its activity of encoding the polypeptide having the specific protease activity; (B) the general tolerance of polynucleotides encoding such proteases to modification and extent of such tolerance; (C) a rational and predictable scheme for modifying any nucleotide on the polynucleotide encoding said protease with an expectation of obtaining the desired biological function; and (D) the specification provides insufficient guidance as to which of the essentially infinite possible choices is likely to be successful .

Thus, applicants have not provided sufficient guidance to enable one of ordinary skill in the art to make and use the claimed invention in a manner reasonably correlated with the scope

Art Unit: 1652

of the claims broadly including polynucleotides with an enormous number of nucleotide modifications to the polynucleotides encoding SEQ ID NOS: 1 or 2. The scope of the claims must bear a reasonable correlation with the scope of enablement (*In re Fisher*, 166 USPQ 19 24 (CCPA 1970)). Without sufficient guidance, determination of polynucleotides having the desired biological characteristics is unpredictable and the experimentation left to those skilled in the art is unnecessarily, and improperly, extensive and undue. See *In re Wands* 858 F.2d 731, 8 USPQ2d 1400 (Fed. Cir, 1988).

Double Patenting

The nonstatutory double patenting rejection is based on a judicially created doctrine grounded in public policy (a policy reflected in the statute) so as to prevent the unjustified or improper timewise extension of the "right to exclude" granted by a patent and to prevent possible harassment by multiple assignees. See *In re Goodman*, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); *In re Longi*, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985); *In re Van Ornum*, 686 F.2d 937, 214 USPQ 761 (CCPA 1982); *In re Vogel*, 422 F.2d 438, 164 USPQ 619 (CCPA 1970);and, *In re Thorington*, 418 F.2d 528, 163 USPQ 644 (CCPA 1969).

A timely filed terminal disclaimer in compliance with 37 CFR 1.321(c) may be used to overcome an actual or provisional rejection based on a nonstatutory double patenting ground provided the conflicting application or patent is shown to be commonly owned with this application. See 37 CFR 1.130(b).

Effective January 1, 1994, a registered attorney or agent of record may sign a terminal disclaimer. A terminal disclaimer signed by the assignee must fully comply with 37 CFR 3.73(b).

Claims 7-34 are rejected under the judicially created doctrine of obviousness-type double patenting as being unpatentable over claims 4-20 of U.S. Patent No. 6,376,227 and claims drawn to "a gene" in copending applications 10/456479, 10/820712, 10/820714, 11/235249,11/318576. An obviousness-type double patenting rejection is appropriate where the conflicting claims are not identical, but an examined application claim is not patentably distinct from the reference claim, because the examined claim is either anticipated by, or would have been obvious over the

Art Unit: 1652

reference claim. See, e.g., *In re Berg*, 140 F.3d 1428, 46 USPQ2d 1226 (Fed. Cir. 1998); *In re Goodman*, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); *In re Longi* 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985). Although the conflicting claims are not identical, they are not patentably distinct from each other. Claims 7-34 of the instant application and claims 4-20 of the reference patent and claims drawn to “a gene” of the copending applications are all directed to polynucleotides encoding polypeptides that have at least 90% sequence identity with SEQ ID NO:1 or 2 (see enclosed sequence alignments). Among all the different polynucleotides claimed in the instant application and in the reference patent and copending applications a number of polynucleotides are identical to one another. The portion of the specification (and the claims) in the reference patent and copending applications that supports the recited polynucleotides includes several embodiments that would anticipate the polynucleotides and the vectors and host cells claimed in claims 7-34 herein. Claims 7-34 of the instant application listed above cannot be considered patentably distinct over claims 4-20 of the reference patent and claims of the copending applications when there is specifically recited embodiment that would anticipate mainly claims 7-34 of the instant application. Alternatively, claims 7-34 cannot be considered patentably distinct over claims 4-20 of the reference patent and claims of the copending applications when there is specifically disclosed embodiment in the reference patent and reference applications that supports claims 4-20 of that patent (and said applications) and falls within the scope of claims 7-34 herein because it would have been obvious to one having ordinary skill in the art to modify claims 4-20 of the reference patent and the claims drawn to a “gene” in the copending applications by selecting a specifically disclosed embodiment that supports those claims. One of ordinary skill in the art would have been motivated to do this

Art Unit: 1652

because that embodiment is disclosed as being a preferred embodiment within claims of the reference patent and applications.

Conclusion

None of the claims are allowable.

Any inquiry concerning this communication or earlier communications from the Examiner should be directed to Manjunath N. Rao, Ph.D. whose telephone number is 571-272-0939. The Examiner can normally be reached on 7.00 a.m. to 3.30 p.m. If attempts to reach the examiner by telephone are unsuccessful, the Examiner's supervisor, Ponnathapura Achutamurthy can be reached on 571-272-0928. The fax phone numbers for the organization where this application or proceeding is assigned is 571-273-8300 for regular communications and for After Final communications. Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the receptionist whose telephone number is 571-272-1600.



Manjunath N. Rao, Ph.D.
Primary Examiner
Art Unit 1652

May 26, 2006

QY 241 TGRNDSMHEAFRGKITLALYALGRTNANDTNGHGHVAGSVLGNKXTNKMAPOANLVF 300
 DB 241 TGRNDSMHEAFRGKITLALYALGRTNANDTNGHGHVAGSVLGNKXTNKMAPOANLVF 300
 QY 301 QSIMDSXGIGLPSNLQTLFSGOAXSAGARIHTNSKGAIVNGAYTTDSRVVDYVRKNDM 360
 DB 301 QSIMDSXGIGLPSNLQTLFSGOAXSAGARIHTNSKGAIVNGAYTTDSRVVDYVRKNDM 360
 QY 361 TILFAAGNEXPNGGTISAPGTAKNAITVGATENLRPSFGSYADININVAQPSRGPFTKG 420
 DB 361 TILFAAGNEXPNGGTISAPGTAKNAITVGATENLRPSFGSYADININVAQPSRGPFTKG 420
 QY 421 RIKPDVMAAGTITLSARSSLAPDSSFWANHDSKAYMGTSMAPIVAGNVAGLRHFVAK 480
 DB 421 RIKPDVMAAGTITLSARSSLAPDSSFWANHDSKAYMGTSMAPIVAGNVAGLRHFVAK 480
 QY 481 NRGITPSPSLKKAALAGAADIGLGYPNGNGMGRVTLDKSLNVAIVNNESSSLSTSOKAT 540
 DB 481 NRGITPSPSLKKAALAGAADIGLGYPNGNGMGRVTLDKSLNVAIVNNESSSLSTSOKAT 540
 QY 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLVLITAPNGTXYVGNDFXXPXXXND 600
 DB 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLVLITAPNGTXYVGNDFXXPXXXND 600
 QY 601 GRNNVENVPINAPQSGTYTIEVOAYNVPGPQFSLAIYN 640
 DB 601 GRNNVENVPINAPQSGTYTIEVOAYNVPGPQFSLAIYN 640

RESULT 2
 US-10-456-479-4
 ; Sequence 4, Application US/10456479
 ; Publication No. US20040072321A1
 ; GENERAL INFORMATION:

APPLICANT: SATO, TOSYOSHI
 APPLICANT: OKUDA, MITSUYOSHI
 APPLICANT: TAKIMURA, YASUSHI
 APPLICANT: SUMITOMO, NOBUYUKI
 APPLICANT: NOMURA, MASAFUMI
 APPLICANT: KOBAYASHI, TOHRU
 TITLE OF INVENTION: ALKALINE PROTEASE
 FILE REFERENCE: 238700US0
 CURRENT APPLICATION NUMBER: US/10/456, 479
 CURRENT FILING DATE: 2003-06-09
 PRIOR APPLICATION NUMBER: JP 2002-186387
 PRIOR FILING DATE: 2002-06-26
 PRIOR APPLICATION NUMBER: JP 2002-304232
 PRIOR FILING DATE: 2002-10-18
 NUMBER OF SEQ ID NOS: 16
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO 4
 LENGTH: 640
 TYPE: PRT
 ORGANISM: Bacillus sp. KSM-KP43
 US-10-456-479-4

Query Match 99.0%; Score 3029; DB 4; Length 640;
 Best Local Similarity 93.3%; Pred. No. 8.7e-261;
 Matches 597; Conservative 0; Mismatches 43; Indels 0; Gaps 0;

QY 1 MRKKKKVFLSVLSAAAILSTVALXNPSAGAKXKXPDLPFGIOTTTDXGFSKQXOGTAA 60
 DB 1 MRKKKKVFLSVLSAAAILSTVALXNPSAGAKXKXPDLPFGIOTTTDXGFSKQXOGTAA 60
 QY 61 FLVESSENVKLXKGLKXKGLFTVPANNKLIHQFNGLIBETKXLEKXTGAULDIYIPYAY 120
 DB 61 FLVESSENVKLXKGLKXKGLFTVPANNKLIHQFNGLIBETKXLEKXTGAULDIYIPYAY 120
 QY 121 IVEYEGDVXKX 180
 DB 121 IVEYEGDVXKX 180

QY 181 GIEIXAOKXSNXNDVXYITAKPEYKXNDVARGIVKADVAQSSYGLYGCGQIYAVADTGID 240
 DB 181 GIEIXAOKXSNXNDVXYITAKPEYKXNDVARGIVKADVAQSSYGLYGCGQIYAVADTGID 240
 QY 241 TGRNDSMHEAFRGKITLALYALGRTNANDTNGHGHVAGSVLGNKXTNKMAPOANLVF 300
 DB 241 TGRNDSMHEAFRGKITLALYALGRTNANDTNGHGHVAGSVLGNKXTNKMAPOANLVF 300
 QY 301 QSIMDSXGIGLPSNLQTLFSGOAXSAGARIHTNSKGAIVNGAYTTDSRVVDYVRKNDM 360
 DB 301 QSIMDSXGIGLPSNLQTLFSGOAXSAGARIHTNSKGAIVNGAYTTDSRVVDYVRKNDM 360
 QY 361 TILFAAGNEXPNGGTISAPGTAKNAITVGATENLRPSFGSYADININVAQPSRGPFTKG 420
 DB 361 TILFAAGNEXPNGGTISAPGTAKNAITVGATENLRPSFGSYADININVAQPSRGPFTKG 420
 QY 421 RIKPDVMAAGTITLSARSSLAPDSSFWANHDSKAYMGTSMAPIVAGNVAGLRHFVAK 480
 DB 421 RIKPDVMAAGTITLSARSSLAPDSSFWANHDSKAYMGTSMAPIVAGNVAGLRHFVAK 480
 QY 481 NRGITPSPSLKKAALAGAADIGLGYPNGNGMGRVTLDKSLNVAIVNNESSSLSTSOKAT 540
 DB 481 NRGITPSPSLKKAALAGAADIGLGYPNGNGMGRVTLDKSLNVAIVNNESSSLSTSOKAT 540
 QY 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLVLITAPNGTXYVGNDFXXPXXXND 600
 DB 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLVLITAPNGTXYVGNDFXXPXXXND 600
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 DB 601 GRNNVENVPINAPQSGTYTIEVOAYNVPGPQFSLAIYN 640

RESULT 3
 US-10-784-870-6
 ; Sequence 6, Application US/10784870
 ; Publication No. US20040142837A1
 ; GENERAL INFORMATION:

APPLICANT: TAKAIWA, MIKIO
 APPLICANT: OKUDA, MITSUYOSHI
 APPLICANT: SAEKI, KATSUHIKA
 APPLICANT: KOBOTA, HIROMI
 APPLICANT: HITOMI, JUN
 APPLICANT: KAGEYAMA, YASUSHI
 APPLICANT: SHIKATA, SHITSUM
 APPLICANT: NOMURA, MASAFUMI
 TITLE OF INVENTION: ALKALINE PROTEASE
 FILE REFERENCE: 0327-0832-0PCT
 CURRENT APPLICATION NUMBER: US/10/784,870
 CURRENT FILING DATE: 2004-02-24
 PRIOR APPLICATION NUMBER: US/09/509,814A
 PRIOR FILING DATE: 2000-04-06
 PRIOR APPLICATION NUMBER: PCT/JP98/04528
 PRIOR FILING DATE: 1998-10-07
 PRIOR APPLICATION NUMBER: JP 9-274570
 PRIOR FILING DATE: 1997-06-08
 NUMBER OF SEQ ID NOS: 24
 SOFTWARE: PatentIn version 3.0
 SEQ ID NO 6
 LENGTH: 640
 TYPE: PRT
 ORGANISM: Bacillus sp.
 US-10-784-870-6

Query Match 99.0%; Score 3029; DB 4; Length 640;
 Best Local Similarity 93.3%; Pred. No. 8.7e-261;
 Matches 597; Conservative 0; Mismatches 43; Indels 0; Gaps 0;

QY 1 MRKKKKVFLSVLSAAAILSTVALXNPSAGAKXKXPDLPFGIOTTTDXGFSKQXOGTAA 60
 DB 1 MRKKKKVFLSVLSAAAILSTVALXNPSAGAKXKXPDLPFGIOTTTDXGFSKQXOGTAA 60
 QY 61 FLVESSENVKLXKGLKXKGLFTVPANNKLIHQFNGLIBETKXLEKXTGAULDIYIPYAY 120
 DB 61 FLVESSENVKLXKGLKXKGLFTVPANNKLIHQFNGLIBETKXLEKXTGAULDIYIPYAY 120

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Db 121 IVEYGDVXSATSTIEHVESVEPYLPYRIDPOLFTKASBLVKVALDTPKONKEVOLR 180
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Db 181 GIEKIAQFALSNDVYITAKPEYKVMNDVARGIVKADVAQSSYGLYGGQGIYAADTGLD 240
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Db 241 TGRNDSMHEAFRGKITALYALGRTNNANDTNGHGTTHVAGSYLGNGXTNKGAPOANLVF 300
Qy 301 QSIINDSGGLGGLPSNLQTLFSGQASAGARIHTNSMGAANVGAATTSRNDVYRKNDM 360
Db 301 QSIINDSGGLGGLPSNLQTLFSGQASAGARIHTNSMGAANVGAATTSRNDVYRKNDM 360
Qy 361 TILPAAGNEKPNCGTISAPGTAKNAITVGATENLRPSFGSYADNINHVAQSSRGPTXOG 420
Db 361 TILPAAGNEKPNCGTISAPGTAKNAITVGATENLRPSFGSYADNINHVAQSSRGPTXOG 420
Qy 421 RIKEDVMAPGTXILSARSSLAPOSSFMANDSKYAMGTSMAPIVAGNVAQLREHFK 480
Db 421 RIKEDVMAPGTXILSARSSLAPOSSFMANDSKYAMGTSMAPIVAGNVAQLREHFK 480
Qy 481 NRGITPKPSLLKALIAAGADXLGYPNGQGMGRVTLDSINVAYNESXLSSTQKAT 540
Db 481 NRGITPKPSLLKALIAAGADXLGYPNGQGMGRVTLDSINVAYNESXLSSTQKAT 540
Qy 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLDLVITAPNGTYVGNDFKXPXXXND 600
Db 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLDLVITAPNGTYVGNDFKXPXXXND 600
Qy 601 GRNNVENVPINXPOSGTYTIEVQAVNVPVGPQPSLAIYN 640
Db 601 GRNNVENVPINXPOSGTYTIEVQAVNVPVGPQPSLAIYN 640

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RESULT 4

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US-10-820-712A-3
; Sequence 3, Application US/10820712A
; Publication No. US20050026804A1
; GENERAL INFORMATION:
; APPLICANT: KAO CORPORATION
; APPLICANT: Okuda, Mitsuyoshi
; APPLICANT: Izawa, Yoshihumi
; APPLICANT: Kobayashi, Tohru
; APPLICANT: Koyama, Shingo
; APPLICANT: Sato, Tsuyoshi
; TITLE OF INVENTION: ALKALINE PROTEASE
; FILE REFERENCE: 251701-USO
; CURRENT APPLICATION NUMBER: US/10/820,712A
; PRIOR FILING DATE: 2003-04-09
; PRIOR APPLICATION NUMBER: 2003-106708
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3
; LENGTH: 640
; TYPE: PRT
; ORGANISM: Bacillus sp. KSM-KP43
US-10-820-712A-3

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Query Match 99.0%; Score 3029; DB 5; Length 640;

Best Local Similarity 93.3%; Pred. No. 8.7e-261; Matches 597; Conservative 0; Mismatches 43; Indels 0; Gaps 0;

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Qy 1 MRKKKKVFLSVLSAAALISTVALNXPAGAXRXPDLDFKGIQTTTDXKXGSKXQOTGAAA 60
Db 1 MRKKKKVFLSVLSAAALISTVALNXPAGAXRXPDLDFKGIQTTTDXKXGSKXQOTGAAA 60

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Qy 61 FLVSEENVKLPKGLQKLETPANNKHLIQFNGPILBETKQLETKAKLIDYIPDYAY 120
Db 61 FLVSEENVKLPKGLQKLETPANNKHLIQFNGPILBETKQLETKAKLIDYIPDYAY 120
Qy 121 IVEYGDVXSXXXIIEHVESVEPYLPXYXIDPOLFTKASXLVKAALDTKONKEVOLR 180
Db 121 IVEYGDVXSATSTIEHVESVEPYLPYRIDPOLFTKASBLVKVALDTPKONKEVOLR 180
Qy 181 GIEKIAQXXXNDVYITAKPEYKVMNDVARGIVKADVAQSSYGLYGGQGIYAADTGLD 240
Db 181 GIEKIAQFALSNDVYITAKPEYKVMNDVARGIVKADVAQSSYGLYGGQGIYAADTGLD 240
Qy 241 TGRNDSMHEAFRGKITALYALGRTNNANDTNGHGTTHVAGSYLGNGXTNKGAPOANLVF 300
Db 241 TGRNDSMHEAFRGKITALYALGRTNNANDTNGHGTTHVAGSYLGNGXTNKGAPOANLVF 300
Qy 301 QSIINDSGGLGGLPSNLQTLFSGQASAGARIHTNSMGAANVGAATTSRNDVYRKNDM 360
Db 301 QSIINDSGGLGGLPSNLQTLFSGQASAGARIHTNSMGAANVGAATTSRNDVYRKNDM 360
Qy 361 TILPAAGNEKPNCGTISAPGTAKNAITVGATENLRPSFGSYADNINHVAQSSRGPTXOG 420
Db 361 TILPAAGNEKPNCGTISAPGTAKNAITVGATENLRPSFGSYADNINHVAQSSRGPTXOG 420
Qy 421 RIKEDVMAPGTXILSARSSLAPOSSFMANDSKYAMGTSMAPIVAGNVAQLREHFK 480
Db 421 RIKEDVMAPGTXILSARSSLAPOSSFMANDSKYAMGTSMAPIVAGNVAQLREHFK 480
Qy 481 NRGITPKPSLLKALIAAGADXLGYPNGQGMGRVTLDSINVAYNESXLSSTQKAT 540
Db 481 NRGITPKPSLLKALIAAGADXLGYPNGQGMGRVTLDSINVAYNESXLSSTQKAT 540
Qy 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLDLVITAPNGTYVGNDFKXPXXXND 600
Db 541 YXFTATAGKPLKISLVMSDAPASTTASVTLVNDLDLVITAPNGTYVGNDFKXPXXXND 600
Qy 601 GRNNVENVPINXPOSGTYTIEVQAVNVPVGPQPSLAIYN 640
Db 601 GRNNVENVPINXPOSGTYTIEVQAVNVPVGPQPSLAIYN 640

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RESULT 5

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US-10-820-714A-3
; Sequence 3, Application US/10820714A
; Publication No. US20050214922A1
; GENERAL INFORMATION:
; APPLICANT: KAO CORPORATION
; APPLICANT: Okuda, Mitsuyoshi
; APPLICANT: Kobayashi, Tohru
; APPLICANT: Sumitomo, Nobuyuki
; APPLICANT: Takimura, Yasuichi
; APPLICANT: Sato, Tsuyoshi
; TITLE OF INVENTION: ALKALINE PROTEASE
; FILE REFERENCE: 251697USO
; CURRENT APPLICATION NUMBER: US/10/820,714A
; PRIOR FILING DATE: 2004-04-09
; PRIOR APPLICATION NUMBER: 2003-106709
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3
; LENGTH: 640
; TYPE: PRT
; ORGANISM: Bacillus sp. KSM-KP43
US-10-820-714A-3

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Query Match 99.0%; Score 3029; DB 5; Length 640;

Best Local Similarity 93.3%; Pred. No. 8.7e-261; Matches 597; Conservative 0; Mismatches 43; Indels 0; Gaps 0;

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Qy 1 MRKKKKVFLSVLSAAALISTVALNXPAGAXRXPDLDFKGIQTTTDXKXGSKXQOTGAAA 60
Db 1 MRKKKKVFLSVLSAAALISTVALNXPAGAXRXPDLDFKGIQTTTDXKXGSKXQOTGAAA 60

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